

<b>05/28/2008 - Wednesday</b>				
<b>Time</b>	<b>Type</b>	<b>Abstract #</b>	<b>Title</b>	<b>Speaker</b>
730 - 830am	Breakfast	x	<b>La Fonda Breakfast Buffet</b> (French "Texas Toast", Fluffy scramble eggs, Grilled breakfast potatoes, Applewood smoked bacon, breads, and fruit, etc.)	x
830 - 845	Intro	x	Welcome Intro	<b>Jose Olivares</b>
<b>845 - 930</b>	<b>Keynote</b>	<b>FF0052</b>	<b>Defining the Human Microbiome: Friends or Family?</b>	<b>Bruce Birren</b>
930 - 1000	Speaker 1	FF0090	Sequence Improvement at the Genome Sequencing Center	<b>Bob Fulton</b>
1000 - 1030	Break	x	<b>Beverages provided</b>	x
1030 - 1100	Speaker 2	FF0125	Metagenomic Approaches Targeted Toward the Human Microbiome	<b>Joe Petrosino</b>
1100 - 1130	Speaker 3	FF0103	Reconstruction of Individual Genomes of Uncultured Microbial Taxa from Complex Communities	<b>Ramunas Stepanauskas</b>
1130 - 1200	Speaker 4	FF0110	Targeted genomics of uncultured microbes: from single cells to populations	<b>Mircea Podar</b>
1200 - 130pm	Lunch	x	<b>Coronado Lunch Buffet</b> (Char-grilled chicken breast with barbecue-chipotle vinaigrette, Pan-seared rainbow trout fillet served with smoked yellow pepper coulis, Roasted garlic mashed potatoes and seasonal vegetables, etc.)	x
130 - 200	Speaker 5	FF0105	New scientific breakthroughs using the 454 Genome Sequencer FLX: 400 base pair reads and more.....Roche 454	<b>Tim Harkin</b>
200 - 230	Speaker 6	FF0102	Next Generation Sequencing - Illuminating the Genome - Illumina corp.	<b>Haley Fiske</b>
230 - 300	Speaker 7	FF0116	High-resolution Structural Variation Detected with Ultra High-throughput Sequencing of Paired End Libraries.....AB SOLiD	<b>Michael Rhodes</b>
300 - 400	<b>Panel Discussion</b>	x	<b>Panel Discussion - Next Generation Sequencing Technologies</b>	Chair - Donna Muzny
400 - 430	Break	x	<b>Beverages and snacks provided</b>	x
415 - 530	<b>Posters - even #s</b>	x	<b>Poster Session</b>	x
530 - 800pm	<b>Meet &amp; Greet Party</b>	x	<b>Meet &amp; Greet Party - sponsored by Roche --- Food &amp; Drinks</b>	x
<b>05/29/2008 - Thursday</b>				
<b>Time</b>	<b>Type</b>	<b>Abstract #</b>	<b>Title</b>	<b>Speaker</b>
730 - 830am	Breakfast	x	<b>Santa Fe Breakfast Buffet</b> (Scrambled eggs with a choice of three accompaniments on the side - chilaquiles with green chile and cheese, chorizo sausage and roasted green chile, Grilled breakfast potatoes, applewood-smoked bacon and warm flour tortillas, assorted breads and fruits, etc.)	x
830 - 845	Intro	x	Welcome Back Intro	<b>Jim Bristow</b>
<b>845 - 930</b>	<b>Keynote</b>	<b>FF0050</b>	<b>The Unfinished Genome</b>	<b>Sydney Brenner</b>
930 - 955	Speaker 1	FF0085	How Close to Finished can an Assembler Get?	<b>Jim Knight</b>
955 - 1020	Break	x	<b>Beverages provided</b>	x
1020 - 1045	Speaker 2	FF0098	Taking the Next Step: Assembling Large Genomes Using NextGen Sequencing	<b>Sean Sykes</b>
1045 - 1110	Speaker 3	FF0034	De Novo Assembly of Microbial Genomes from Illumina Whole-Genome Shotgun Data	<b>Iain MacCallum</b>
1110 - 1135	Speaker 4	FF0036	Combining Next-Gen Sequence Technologies in Multi-platform Assemblies	<b>Christian Buhay</b>
1135 - 1200	Speaker 5	FF0010	The Arcturus - Minerva assembly management system	<b>Ed Zuiderwijk</b>
1230 - 130pm	Lunch	x	<b>New Mexican Lunch Buffet</b> (Pork tenderloin achiote-rubbed and char-grilled with tomatillo-chipotle sauce, your choice of either Chicken or Cheese enchiladas with red or green chile, etc.)	x
130 - 155	Speaker 6	FF0076	Bacterial Genome Finishing on the Turn of New Technology	<b>Hajnalka Kiss</b>
155 - 220	Speaker 7	FF0088	Towards "Finishing" of Eukaryotic Transcriptomes	<b>Stephen Kingsmore</b>
220 - 245	Speaker 8	FF0075	An optimized DNA sequencing pipeline to support drug discovery at Wyeth	<b>Jan Kieleczawa</b>
245 - 310	Speaker 9	FF0065	Sequencing the Genome, Transcriptome, Methylome and smRNAome of the Arabidopsis Ecotype, Cape Verde Island	<b>Ronan O'Malley</b>
310 - 415	<b>Panel Discussion</b>	x	<b>Panel Discussion - Sequencing and Finishing Standards</b>	Chair - Darren Grafham
415 - 630	<b>Wine &amp; Cheese</b>	x	<b>Beverages, Wine &amp; Cheese provided - sponsored by Applied Biosystems</b>	x
415 - 630	<b>Posters - odd #s</b>	x	<b>Poster Session with Wine &amp; Cheese - sponsored by Applied Biosystems</b>	x
630 - bedtime	on your own	x	Dinner and night on your own - enjoy	x
<b>05/30/2008 - Friday</b>				
<b>Time</b>	<b>Type</b>	<b>Abstract #</b>	<b>Title</b>	<b>Speaker</b>
730 - 830am	Breakfast	x	<b>Healthy Start Breakfast Buffet</b> (Scrambled Eggs on side tomatoes, scallions and spinach, Turkey sausage links, Assorted chilled fruit juices, Platter of freshly sliced seasonal fruit, Assorted and bran muffins with butter, Granola and oatmeal served with low-fat milk, Individual assorted fruit yogurts, etc.)	x
830 - 845	Intro	x	Welcome Back Intro	<b>Chris Detter</b>
<b>845 - 930</b>	<b>Keynote</b>	<b>FF0053</b>	<b>Great Expectations: Fulfilling the promises of the Human Genome Project</b>	<b>Deanna Church</b>
930 - 1000	Speaker 1	FF0087	Using Consed and Cross_match in Resequencing Projects	<b>David Gordon</b>
1000 - 1030	Break	x	<b>Beverages and snacks provided</b>	x
1030 - 1100	Speaker 2	FF0008	1000 Genomes Project Data Management and Analysis	<b>Hoda Khouri</b>
1100 - 1130	Speaker 3	FF0020	Genome annotation improvement using new massive sequencing technologies.	<b>François Artiguenave</b>
1130 - 1200	Speaker 4	FF0041	AlpheusTM - a system for nucleotide variant detection and digital gene expression analysis in ultra-high throughput sequencing projects	<b>Neil Miller</b>
1200 - 1230	<b>Closing Discussions</b>	x	<b>Closing Discussions - discuss next year's plans</b>	Chair - Chris Detter
1230 - 200pm	Lunch & Close of meeting	x	<b>La Fiesta Plaza Lunch Buffet</b> - (Chicken and beef fajitas with grilled red onions and bell peppers, Black beans (Vegetarian), Spanish rice (Vegetarian), Pork posole and calabacitas rancheras, Warm flour tortillas and butter, etc.) <b>End of meeting, enjoy lunch and Santa Fe</b>	x